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; Search time 194.042 Seconds (without alignments) 10829.323 Million cell updates/sec
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Listing first 45 summaries
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1410
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

282 ATGCCCTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGA 341

222 TGCTGGTGCTTAACCAAATCCACCTGCTACTGCTAATTTAGTCACATAATGTAAGGTTAA 281 

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342 AIGTGTTAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGC 401

255 TGCTACAGCTACTGCTACTGCTACTGCTACTGCTACTGCTACTGCTACTGCTAC 314

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435 TGCTACTGCTACTGCTACTGCTACTGCTACTGCTACTGCTACTGCTACTGCTAC 494

522 AGTAACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACTTAACTTTA

CTATAATGGTAATAGTAATACTCCTTTCAATCCAGGTAAAAGTTAATGCACACCTTG

	Description		Socialize 3938, A			Sequence 3500, Ap	4034	1140,	1170	22, Ap	1913,	4707,	Sequence 7632, Ap	Sequence 2529, Ap		22	100	Segmence 20125	10		5 0	2047	Z047,	Seguence 119, App	48328	8573	, ,	Seguence 1, Applit	۲	,
SUMMARIES	ID	US-10-144-771-39592	US-10-144-771-31796	US-10-203-1388-8641	0-203-1	2 0	IIS-10-203-1387-7146	10-002-110-01	17-804-708-91		-10-144-771-470	10-144-7/1-4/U/	10 203 -138A-7	US-10-203-138A-2529	US-10-194-046-1	US-10-092-411A-2243	US-09-949-004-601	US-10-144-771-29725	US-10-311-506-48	IIS-10-311-507-48	US-10-017-161-2047	115-10-017-161-2040	****	02-03-348-T74-TTA	US-09-531-113-48328	US-10-203-138A-8573	US-09-692-570-1	US-10-144-771-22890	5-09-948-12	
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227, A e 1, Ap 6082,	Sequence 1937, Ap Sequence 2000, Ap Sequence 31479, A Sequence 22, Appl. Sequence 2103, Appl. Sequence 2113, A Sequence 2113, A Sequence 294, Appl. Sequence 7446, Ap Sequence 7446, Ap Sequence 1, Appl. Sequence 5026, Appl. Sequence 5036, Ap	O USES THEREOF	Length 897; Indels 0; Gaps 0;
574 3673778 756 9180	19180 6 17594 6 729 6 1118 6 1118 6 11083 5 1372 6 5912 6 6092 6 74105 5 1278 6 1278 6	MENTS VERY SYSTEM A	5.5%; Score 77.4; DB 6; L. 43.9%; Pred. No. 4.3e-08; vative 0; Mismatches 421;
42 42 41.8 81.8	11111111111111111111111111111111111111	RESULT 1 US-10-144-771-39592 Sequence 39592, Application US/10144771 GENERAL INFORMATION: APPLICANT: VENTER, J. Craig TITLE OF INVENTION: HUMAN GENOME DISCO FILE REPERENCE: CLOOL321 CURRENT FILING DATE: 2002-05-15 NUMBER OF SEQ ID NOS: 47235 SEQ ID NO 39592 LENGTH: 897 TYPE: DNA OCCANISM: HUMAN US-10-144-771-39592	Query Match Best Local Similarity Matches 330; Conser

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ORGANISM: Homo sapiens
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GCT 206
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Pred. No. 2.1e-05;
0; Mismatches 411;
                                                                                                                                                                                                                                                                                                      Sequence 3176, Application US/10144771
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig
TITLE OF INVENTION: HUMAN GENOME DISCOVERY SYS
FITLE REFERENCE: CLO01321
CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 47235
SEQ ID NO 31796
LENGTH: 961
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Matches 312; Conservative
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ORGANISM: HUMAN
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APPLICANT: Molecular Lyndamics, auc.
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, David R.
APPLICANT: Hanzel, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUWAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USFFUL FOR TITLE OF INVENTION: HUWAN GENOME-02
TITLE OF INVENTION: HUWAN GENOME-02
TITLE OF INVENTION: HUWBER: US/10/203,138A
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: OF MAY 2000 (04.02.00)
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/234,366
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: OF September 2000 (03.10.00)
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: US September 2000 (21.09.00)
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,4687
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APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
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Thu Feb

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TCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAAATGCCCTGCTGGTACCGC 299
                                                                                                                                   300 AATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGTTAATTGTAGAAT 359
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RESULT

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GENERAL INNCHMANTION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USPENDED STORM TO HUMAN GENOME CONTREMY APPLICATION NUMBER: US/10/203,138A
CURRENT FILING DATE: 2002-080-02
PRIOR APPLICATION NUMBER: US 60/203,436
PRIOR APPLICATION NUMBER: US 60/203,436
PRIOR FILING DATE: 20 May 2000 (26.05.00)
PRIOR FILING DATE: 20 May 2000 (26.05.00)
PRIOR FILING DATE: 30 August 2000 (33.08.00)
PRIOR FILING DATE: 21 September 2000 (31.09.00)
PRIOR FILING DATE: 21 September 2000 (27.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 21 September 2000 (31.09.00)
PRIOR FILING DATE: 21 September 2000 (31.09.00)
PRIOR FILING DATE: 30 August 2000 (30.06.00)
PRIOR FILING DATE: 21 September 2000 (31.09.00)
PRIOR FILING DATE: 30 September 2000 (30.06.00)
PRIOR FILING DATE: 30 SEPTEMBER OF SEQ ID NOS: 15438
SOFTWARE: MOISCULAR DYNAMICS SEQUENCE LISTING ENGLISH: 1973
"TUPE: NA
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4.3%; Score 60; DB 6; Le
Best Local Similarity 41.5%; Pred. No. 0.00037;
Matches 463; Conservative 0; Mismatches 650;
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Sequence 3500, Application US/10203138A GENERAL INFORMATION:
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TAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTGAAGCCACTGCAGG 896
                                                                                                                                                                                                                                                                                                TGGTGCCGCTACTTTAGCCAAATAATGTAATATTGCATGCCCTGATGGTACTGCAATTGC 956
                                                                                                                                                                                                                                                                                                               TGTTAGATCATTCACAGAATGTGATTAAATGTAGACTTAACTTTTACTATAATGGTAATAA 596
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GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig
TITLE OF INVENTION: HOWAN GENOME DISCOVERY SYSTEM AND USES THEREOF
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CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
NUMBER OF ESQ ID NOS: 47235
SEQ ID NO 4034
LENGTH: 1188
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US-10-144-771-4034/c
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CRGANISM: HUMAN
US-10-144-771-4034
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APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
FILE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: OF MAGUET 2000 (04.02.00)
PRIOR FILING DATE: OF MAGUET 2000 (03.08.00)
PRIOR FILING DATE: 03 AUGUST 2000 (03.08.00)
PRIOR FILING DATE: 03 OCTCOBER 2000 (03.08.00)
PRIOR FILING DATE: 03 OCTCOBER 2000 (03.08.00)
PRIOR PRILING DATE: 03 OCTCOBER 2000 (03.08.00)
PRIOR FILING DATE: 03 OCTCOBER 2000 (03.08.00)
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PRIOR FILLING DATE: 21 September 2000 (27.09.00)
PRIOR FILLING DATE: 21 September 2000 (21.09.00)
PRIOR PPLICATION NUMBER: US 09/608,408
PRIOR FILLING DATE: 30 June 2000 (30.06.00)
PRIOR FILLING DATE: 15438
SOFTWARE: Molecular Dynamics Sequence Listing Engine
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US-10-203-138A-7146/C
US-quence 7146, Application US/10203138A
; GENERAL INFORMATION:
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GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-101
CURRENT FILING DATE: 2002-03-07
FRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1997-08-14
PRIOR FILING DATE: 1997-08-14
NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER: US 60/055,779
SEQ ID NO 1120
LENGTH: 1248
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3.7%; Score 52.2; DB 6; Length 510;
Best Local Similarity 46.2%; Pred. No. 0.017;
Matches 174; Conservative 0; Mismatches 203; Indels
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3.5%; Score 50; DB 6; Length 1248;
Best Local Similarity 42.8%; Pred. No. 0.06;
Matches 254; Conservative 0; Mismatches 340; Indels
                                                                                                                                     ; OTHER INFORMATION: NT HIT: U01287.1, EVALUE 1.30e+00 US-10-203-138A-7146
                                                                                                    OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 4.6
                                                                      OTHER INFORMATION: MAP TO AL035457.11
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             LENGTH: 510
TYPE: DNA
ORGANISM: Homo sapiens
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SEQ ID NO 7146
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; LOCATION: (1)..(1141)
; OTHER INFORMATION: consensus sequence of A.t., L.a., and B.n. FAE1 promoters
US-09-806-708B-22
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GCTTGTGCTACGGGGACTAATTCAATAGGTGAGGCATTTAAAATTATTCAACGTGGTAT 552
                                                                                                                                                              TATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTACATGCACACGCTTGTCCGGTA 426
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GENERAL INFORMATION:
APPLICANT: The University of British Columbia
TILLE OF INVENTION: Regulation of Embryonic Transcription in Plants
FILE REFERENCE: 4810-5841
CURRENT APPLICATION NUMBER: US/09/806,708B
CURRENT FILING DATE: 2001-04-03
PRIOR FILING DATE: 1999-08-04
NUMBER OF SEQ ID NOS: 23
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3.5%; Score 49; DB 5; Length 114
Best Local Similarity 11.5%; Pred. No. 0.099;
Matches 121; Conservative 365; Mismatches 557; Indels
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SEQ ID NO 22
LENGTH: 1141
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ATAAAGTTTAAGGCGCTGTAGCAACTGCAGGTGGTACTGCTACTTTAATTGCATAATGTG 1135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGCAACTAATTATGTAATATTATAAACAGAATGTCTAAATTG-----TGCTGCTAACT 1015
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                                                                                                                           HRRRYNNGWIBAMAYRRWIMNNNNNAKAMCKRAKYWGWNRABVNST-CITWKSKTIKVR 317
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139 HHYVTAMNNAWTTMCMMDKDDKRTRWWWKKNNNATGWDDDTKYHWWNNGCBTVTWMVRY 198
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DDRWRBAYTNNNNNRAXYGAYADDYAYYMSDTCDAWMKWDATKMNNATTYNRGTAWRT
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                                                                                                                                                                      423 GGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAATGCCGCTACCATAGTCGCATAATG
                                  303 TGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGTTAATTGTAGAATTAA
                                                           363 TITITATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTACATGCACAGCTTGTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SUWA, MAKIKO APPLICANT: ASAI, KIYOSHI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        513 TGATGATGGAGTAACTACTGATTATGTTAGATCATTCACAGAATGTGTTAAATGTAGACT 572
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TITLE OF INVENTION: HUMAN GENOME DISCOVERY SYSTEM AND USES THEREOF
FILE REFERENCE: CL001321
CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 47235
SEQ ID NO 4707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6; Length 3532;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
3.4%; Score 48; DB 6; Length 353
Best Local Similarity 44.0%; Pred. No. 0.19;
Matches 204; Conservative 0; Mismatches 260; Indels
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                                              RECEPTORS
APPLICANT: AKIYAMA, YUTAKA
APPLICANT: ABUBATANI, HIROYUKI
TILE OF INVENTION: NOVEL G PROTEIN-COUPLED REC
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT APPLICATION NUMBER: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTHARE: PATENTIN VET: 2.1
SEQ ID NO 1913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-144-771-4707/c; Sequence 4707, Application US/10144771; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                ; NAME/KEY: CDS
; LOCATION: (2622)..(3332)
US-10-017-161-1913
                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: CDS
LOCATION: (201)..(437)
                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                             LOCATION: (1)..(3532)
                                                                                                                                                                                                                                                                                                                            NAME/KEY: source
                                                                                                                                                                                                                                                              TYPE: DNA
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APPLICANT: MOLECULAR Dynamics, Inc.
APPLICANT: MOLECULAR Dynamics, Inc.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
FILE REFERENCE: PB 0004 WO 8
CURRENT APPLICATION NUMBER: US/10/203,138A
CURRENT APPLICATION NUMBER: US 60/180,312
                                                                                                                                                                                                                              348 TAATTGTAGAATTAATTTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTAC 407
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                                                                                                                      Length 1075;
                  EST_HUMAN HIT: AV739739.1, EVALUE 1.00e+00
                                                                                                              Score 45.8; DB 6; Length 1:
Pred. No. 0.51;
0; Mismatches 402; Indels
                                                    ; OTHER INFORMATION: NT HIT: AL163201.2, EVALUE 2.00e-19 US-10-203-138A-7632
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                                                                                                              3.28;
                                                                                                                                   Best Local Similaria,
Matches 287; Conservative
                                                                                                                                 Similarity
              OTHER INFORMATION: FEATURE:
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US-10-203-138A-2529
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GENERAL INFORMATION:
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Melecular Dynamics, Inc.
APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUARN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUARN GENOME-DERIVED SINGLE EXPRESSION IN BT 474
TITLE OF INVENTION: HUARN GENOME-DERIVED SINGLE EXPRESSION IN BT 474
TITLE OF INVENTION: HUARN GENOME-DERIVED SINGLE EXPRESSION IN BT 474
TITLE OF INVENTION: HUARN GENOME-DERIVED SINGLE EXPRESSION IN BT 474

CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 03 Adust 2000 (04.02.00)
PRIOR FILING DATE: 03 Adust 2000 (04.02.00)
PRIOR FILING DATE: 03 Adust 2000 (03.08.00)
PRIOR FILING DATE: 03 Ctober 2000 (03.10.00)
PRIOR FILING DATE: 103 Cotober 2000 (03.10.00)
PRIOR FILING DATE: 103 Cotober 2000 (03.10.00)
PRIOR FILING DATE: 21 September 2000 (27.09.00)
PRIOR FILING DATE: 31 September 2000 (21.09.00)
PRIOR FILING DATE: 30 June 2000 (30.00)
PRIOR FILING DATE: 31 September 2000 (30.00)
PRIOR FILING DATE: 31 September 2000 (30.00)
PRIOR FILING DATE: 30 June 2000 (30.00)
PRIOR FILING DATE: 30 June 2000 (30.00)
PRIOR FILING DATE: 30 June 2000 (30.00)
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                                                                                                                                                                                                                                                                                400 GCTAGTACATGCACAGCTTGTCCGGTAAACAGAGTTGGTGGTGCATTGACTGCTGGTAAT 459
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                                                                                                 Score 47.4; DB 6; Length 1
Pred. No. 0.24;
0; Mismatches 171; Indels
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LENGTH: 1075
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OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL -
FEATURE:
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                                                                                               3.4%;
                                                                                         Query Match 3.45
Best Local Similarity 46.75
Matches 150; Conservative
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ORGANISM: Homo sapiens
                  ; TYPE: DNA
; ORGANISM: HUMAN
US-10-144-771-4707
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US-10-203-138A-7632
LENGTH: 1954
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PRIOR FILING DATE: 04 February 2000 (04.02.00)
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 26 May 2000 (26.05.00)
PRIOR FILING DATE: 30 August 2000 (33.08.00)
PRIOR FILING DATE: 03 August 2000 (03.08.00)
PRIOR FILING DATE: 03 OCTOBER 2000 (03.10.00)
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 27 September 2000 (27.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 30 September 2000 (21.09.00)
PRIOR FILING DATE: 31 September 2000 (21.09.00)
PRIOR FILING DATE: 30 June 2000 (30.06.00)
NUMBER OF SEQ ID NOS: 15438
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Pred. No. 0.53;
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                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: MAP TO AL078472.1
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Local Similarity 41.7%;
Nes 287; Conservative C
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                          SEQ ID NO 2529
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Matches
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Sequence 2243, Application US/10092411A
Sequence 2243, Application US/10092411A
Sequence 2243, Application US/10092411A
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCC
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-101
CURRENT APPLICATION NUMBER: US/10/092,411A
CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: US 09/134,001
PRIOR FILING DATE: 1998-08-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                            APPLICANT: Robertson, Erle S.
TITLE OF INVENTION: LANA Viral Protein Binding Sites FILE BEFRENCE: UM-07142
CURRENT APPLICATION NUMBER: US/10/194,046
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: 09/410,399
PRIOR FILING DATE: 1999-10-01
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 44.0%; Pred. No. 0.6;
Matches 194; Conservative 0; Mismatches 247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Kaposi's sarcoma-associated herpesvirus
                       1226 TGGTGGTGGTTGTGGTAGTGGTGGTG 1254
948 IGCAATIGCTAGIGGAGCAACTAATTAIG
                                                                                                                                                                     Sequence 1, Application US/10194046 GENERAL INFORMATION:
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US-10-092-411A-2243
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Sequence 601, Application US/09949004
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN HUMAN G-PROTEIN COUPLED
TITLE OF INVENTION: RECEPTOR GENES ASSOCIATED WITH DISEASE, METHODS OF DETECTION
TITLE OF INVENTION: AND USES THEREOF
TITLE OF INVENTION: AND USES THEREOF
TITLE OF INVENTION: AND USES THEREOF
CURRENT FILING DATE: 2002-12-20
PRIOR FILING DATE: 2000-09-13
NUMBER OF SEQ ID NOS: 6961
SEQ ID NOS: 6961
SEQ ID NO 601
LENGTH: 21423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             498 TACTGGTACTGCACTTGATGATGAGTAACTACTGATTATGTTAGATCATTCACAGAATG 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                846 ATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGCTGAAGCCACTGCAGGTGGTGCCGC
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                                                                                                                                                                                                                            DB 6; Length 11091;
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                                                                                                                                                                                                                 Query Match
3.2%; Score 45.8; DB 6; Length 1
Best Local Similarity 46.9%; Pred. No. 0.71;
Matches 143; Conservative 0; Mismatches 162; Indels
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0; Mismatches 290;
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
LENGTH: 11091
                                                                                                                                                           ; ORGANISM: Staphylococcus epidermidis US-10-092-411A-2243
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                                                                                                                                     AAGTGCTGCTGGAGTAAATTGGGTAGCACAAAACACTGAATGTACTAATTGTGCTCC 797
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                                                                  TTTAGGTAATGATGCTACAATAACCGCATAATGTAACGTTGCATGCCCTGATGGTACTAT 737
618 AGGTAAAAGTTAATGCACACCTTGTCCGGCAATTAAACCTGCTAATGTTGCTTAAGCTAC
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